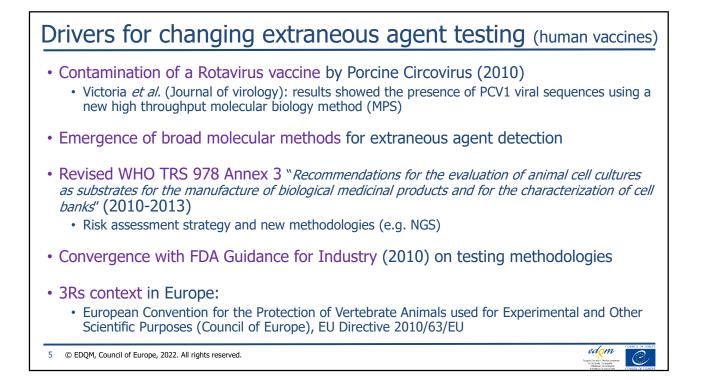
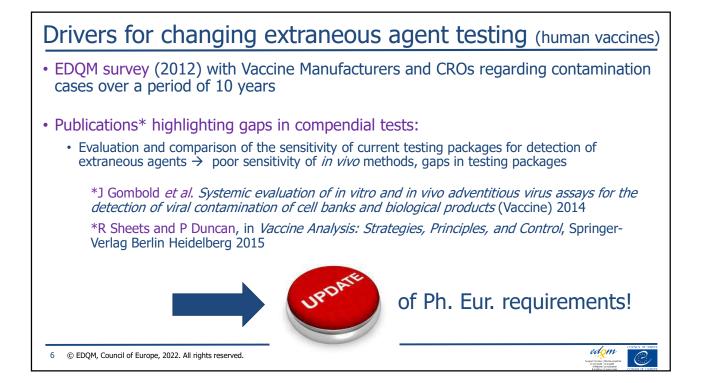


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Outline	
Extraneous agent testing for vaccines: evolution of the Ph. Eur.	
 Vaccines for human use Drivers for revising Ph. Eur. requirements Evolution of Ph. Eur. 5.2.3 & 2.6.16 The concept of Substitution to replace <i>in vivo</i> methods as described in Ph. Eur. 5.2.14 	
 Vaccines for veterinary use Drivers for revising Ph. Eur. requirements Evolution of Ph. Eur. chapters for veterinary vaccines New Approach brings opportunities & benefits Support to stakeholders in a nutshell 	
 High Throughput Sequencing for the detection of extraneous agents 	
► Conclusion	
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	01/2018:50203	• 07/:	2020:20616	01/2018:50214
2.3. CELL SUBST RODUCTION OF IUMAN USE	rrates for the f vaccines for with diploid cell lines and must lines and must lines and	6. TESTS FOR EXTRANEOU ENTS IN VIRAL VACCINES F MAN USE DDUCTION egy for testing extraneous agents in viral va e developed based on a risk assessment foll les of viral contamination risk detailed in p	FOR THE QI VACCINES PURPOSE	STITUTION OF IN VIVO) BY IN VITRO METHOD(S) UALITY CONTROL OF
	Ph. Eur. Chapter 5.2.3 Cell substrates for the production of vaccines for human use	Ph. Eur. Chapter 2.6.16 Tests for extraneous agents in viral vaccines for human use	Ph. Eur. Chapter 5.7 Substitution of in vi methods for the QC vaccines	VO of of of of of of of of of of
Scope	Testing of cell substrates (including extraneous agent testing)	Extraneous agent testing of viral seed lots/harvests	Concept of Substitu to replace <i>in vivo</i> methods	riples described may also ra.
Year introduced or year of last major update	July 2017 (Ph. Eur. Suppl. 9.3)	July 2017 (Ph. Eur. Suppl. 9.3)	July 2017 (Ph. Eur. Suppl. 9.3)	 Revision of chapters 5.2.3 (2.6.16) Elaboration of chapter 5.2.1 (concept of Substitution)

5.2.3 Cell substrates for the production of vaccines for human use

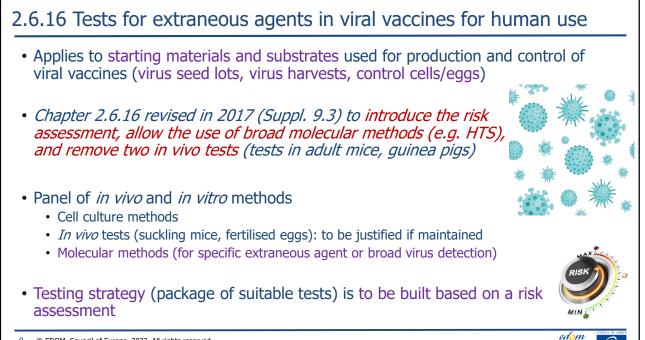
- Scope: diploid cell lines and continuous cell lines used as cell substrates for the production of vaccines
- Chapter 5.2.3 revised in 2017 (Suppl. 9.3) to introduce the risk assessment, allow the use of broad molecular methods (e.g. HTS), and remove an in vivo test (test in adult mice)



- <u>Extraneous agents</u>: testing strategy is to be based on a risk assessment considering e.g. choice of permissive cells, nature of cell lines (e.g. insect cells), cell lines shown to express endogenous retroviral particles, *in vivo* tests to be justified if maintained
- A strategy is given in chapter 5.2.3. Alternative strategies could focus on more extensive testing of the MCB or WCB

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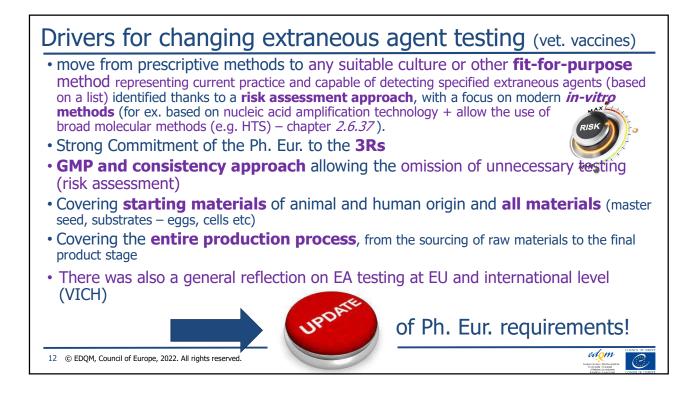




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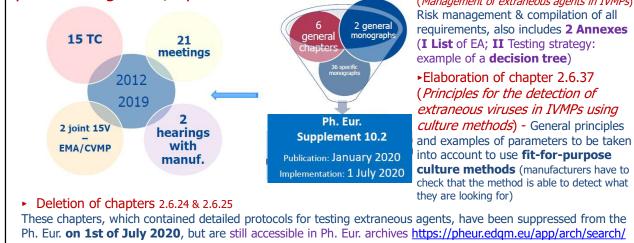
5.2.14 Substitution of <i>in vivo</i> methods for the QC of vaccines
 Chapter elaborated to facilitate the transition to <i>in vitro</i> methods (e.g. HTS), applies to human and vet vaccines
• Chapter 5.2.14 provides guidance on how to introduce alternative <i>in vitro</i> methods, where a head-to-head comparison is not possible
 Envisages the possibility that the relevance and performance of the <i>in vitro</i> method be demonstrated without such head-to-head comparison: concept of "substitution" as an alternative approach for replacement
 Focus on the scientific rationale behind the <i>in vitro</i> methods and the validation package
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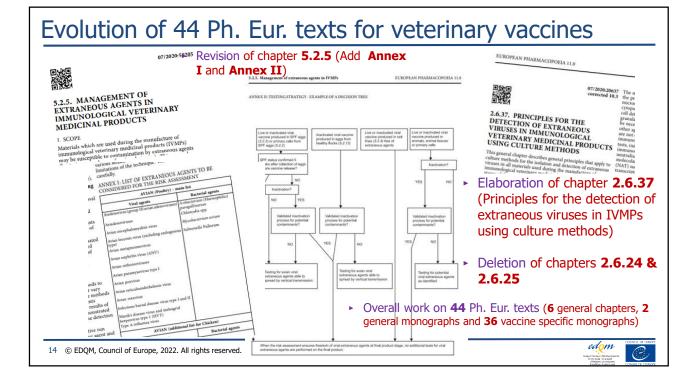


Evolution of Ph. Eur. chapters for veterinary vaccines

Work with all stakeholders: hearings, collaboration with international partners e.g. EMA, April 2020 webinar • Revision of chapter 5.2.5 (Management of extraneous agents in IVMPs)



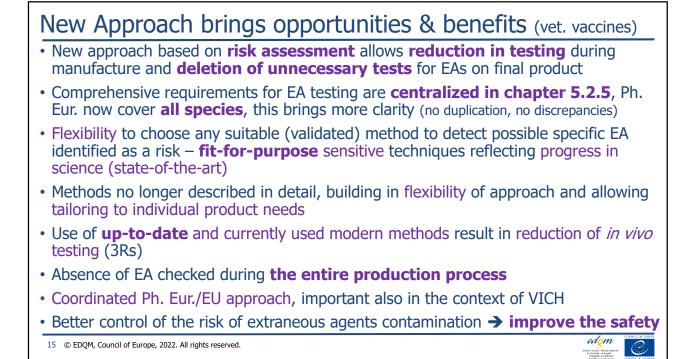
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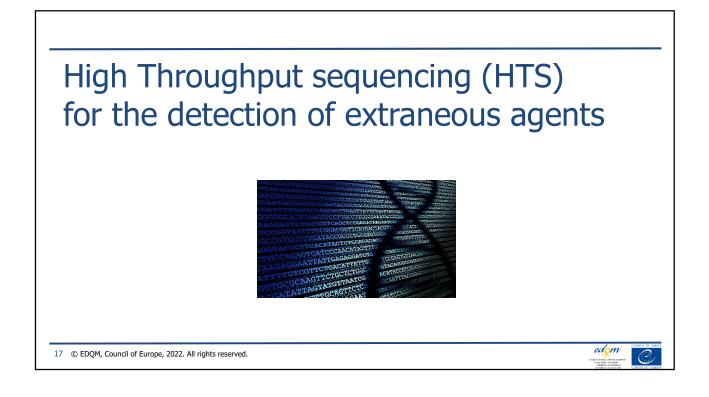
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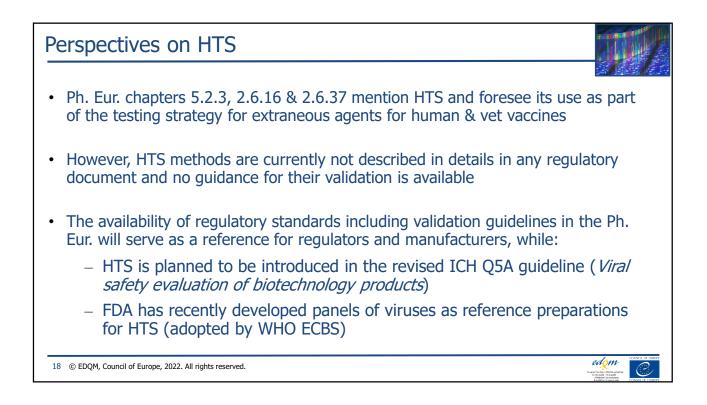
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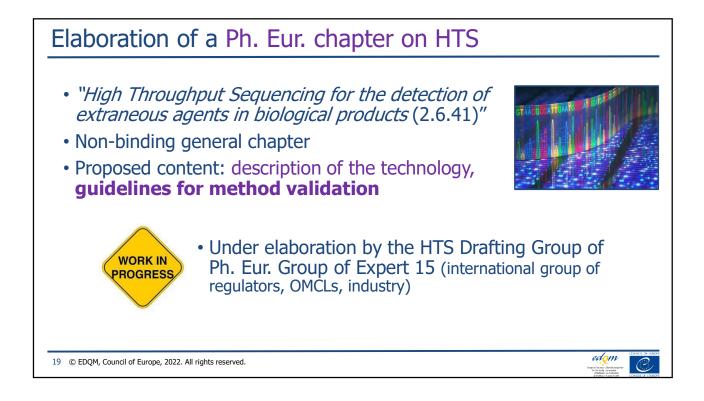
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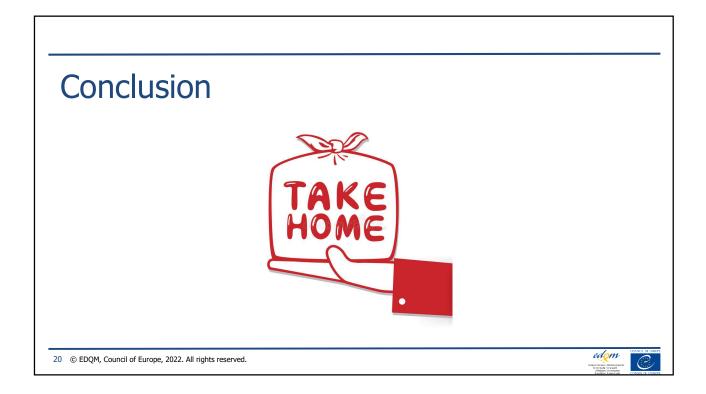


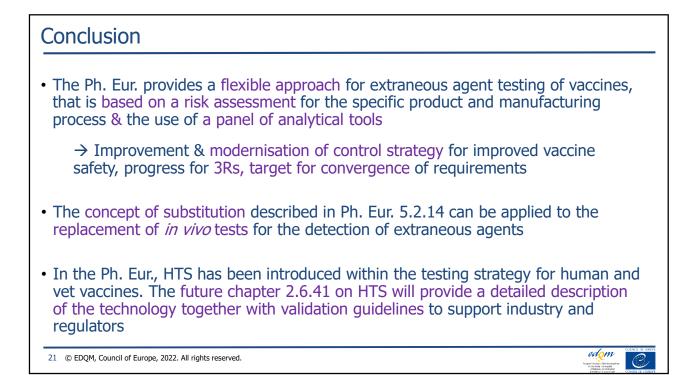
















High Throughput Sequencing: How could the Ph. Eur. help in the exercise to validate HTS methods?

Siemon Ng

Collaboration, Innovation and Scientific Excellence: the European Pharmacopoeia 11th Edition Date: 19-21 September 2022

Outline

- High Throughput Sequencing Technology
- Adventitious Virus Detection in Biologics
- Recent Advancements and Progress in HTS adventitious agent detection
- A New Ph. Eur General Chapter For HTS
- Development in Other Areas
- Summary and Conclusions

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High Throughput Sequencing Technology

• High-throughput sequencing (HTS) are technologies that can rapidly sequence millions of DNA (or RNA) from a single biological samples. HTS has quickly become an important tool in scientific research, drug development and medical diagnostics.

 Multiple technologies and platforms Read length from a hundreds of nucleotides to 50+ Kb Short-reads (e.g. Illumina, Ion Torrent) Long-reads technology (e.g. PacBio, Oxford Nanopore) Up to 10 billion reads simultaneously Direct sequencing of DNA and RNA molecules Epigenetics and modified bases 	Sar	
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Application of HTS as an Analytical Platform

- <u>Adventitious agent detection:</u> Detect and identify both known and unknown adventitious agents with a high level of sensitivity.
- <u>Assessment of genomic stability or variants</u>: Detect low frequency variants or genomic rearrangements
- <u>Identity testing:</u> Verification of engineered construct(s) or mutations at multiple loci simultaneously.
- <u>Genome characterization:</u> Whole genome sequencing to obtain baseline genomic information

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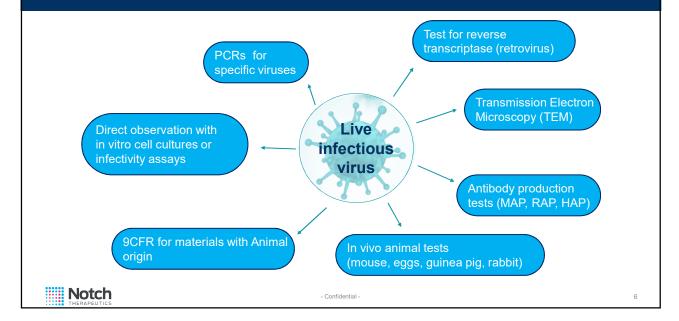
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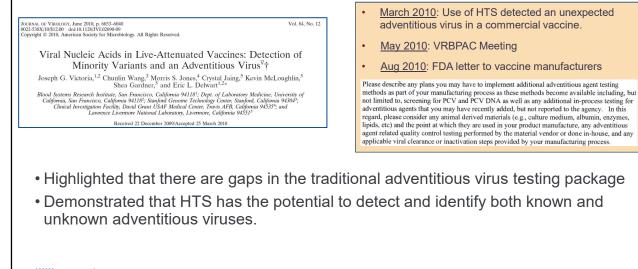
Adventitious Virus Detection In Biologics

Viral Safety Testing Package and Adventitious Virus Testing

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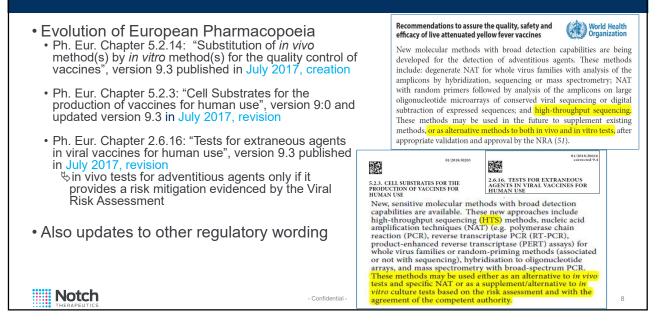
HTS as an Emerging Technology

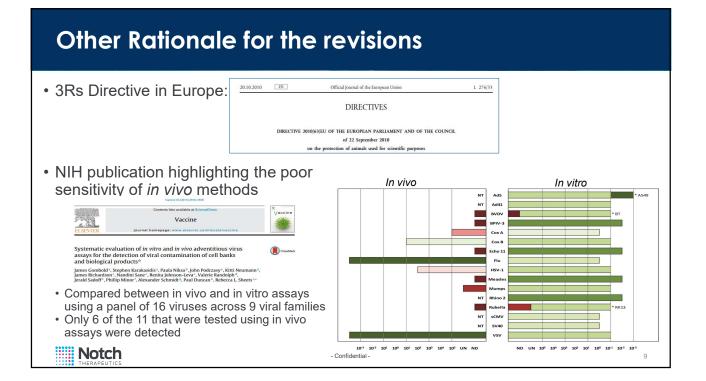


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Updates to European Phamacopoeia to include HTS

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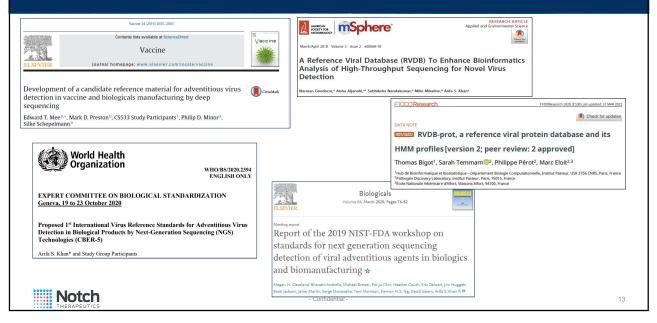
Scientific Advancements

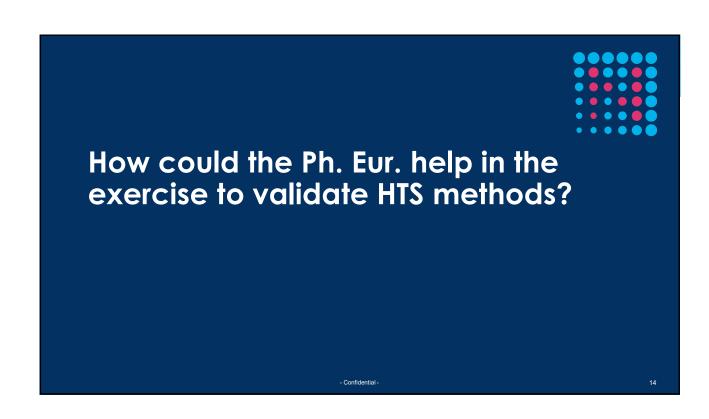
Biologicals 67 (2020) 94-111 Contents lists available at Sciences Biologicals			Biologicals ELSEVIER Volume 59, May 2019, Pages 29-36
ELSEVIER journal homepage: www.elsevier.com/los	cate/biologicals		Use of a new RNA next generation sequencing
Report of the second international conference on ne sequencing for adventitious virus detection in biolog animals®	tics for humans and		approach for the specific detection of virus infection in cells Autre from ". Genin Graid". Enin Marth ". Budof Zirwan ". Januar Chent ". Charles Hoter". Jans-Marie
Arifa S. Khan ^{5,*} , Johannes Blümel ^b , Dieter Deforce ^c , Marion F. Gru Ivana Knezevic ⁶ , Laurent Mallet ^{f,1} , David Mackay ⁸ , Jelle Matthijns			Charpin ⁶ , Alice Marinaci ^b , Benoit Flan [®] , Horst Ruppach [®] , Pascale Beurdeley ⁶ , Marc Eloit ^{6, 6} , ⁶ , ⁶ , ⁶
Sebastiaan Theuns ^e , Joseph Victoria [*] , Pieter Neels ^{el}	🏶 viruses	MDP	
npj Vaccines	Protocol LABRADOR—A Computa in High-Throughput Seque	tional Workflow for Virus Detection encing Data	Approach for the Detection of Viral Contaminants in
	Izabela Fabiańska *, Stefan Borutzki, Benjamin	Richter, Hon Q. Tran, Andreas Neubert and Dietmar Mayer *	Biopharmaceutical and Vaccine Manufacturing Applications Using Next-Generation Sequencing
ARTICLE OPEN	Check for updates		Madolyn L, MacDonald an Shawn W, Polson Md Relyin H, Leena
Sensitivity and breadth of detection			
sequencing for adventitious virus d	etection		RESEARCH ARTICLE
Robert L. Charlebois ¹ , Sarmitha Sathiamoorthy ² , Carine Logvinoff ² , Lucy Gi			
	npj Vaccines	www.nature.com/npjvaccines	Civita for appleter
		Corrected: Author correction	
	ARTICLE OPEN	of an efficient method for the	A Multicenter Study To Evaluate the Performance of High-Throughput
	Server and enterer and the	acids from complex biologicals	Sequencing for Virus Detection
	Sarmitha Sathiamoorthy ^{1,2} , Rebecca J. Malott ¹ , Lucy G		Artfa S, Khan,* Siemon H, S, Ng.* Olivier Vandeputte,< Alsha Allanahl.**
Notch	same same only , needed , Malott , Eddy d	- Confidential -	Avisek Deyati, ⁴ Jean-Pol Cassart, ^a Robert L. Charlebols, ^b Lanyn P. Taltaferro**
THERAPEUTICS		Confidential -	11

Alignment in perspectives

Journal of Chinal Virology 135 (2023) 19412 General line available at Inducedirest Journal of Clinical Virology ELSEVIER journal homepage, www.elsevier.com/houted/pv	Jaural of China' Visiog 154 (2021) 194084 Control for stability of China' Distort Journal of China' Distort ELSIVER journal homegage, www.absorber.com/chantyor	PDA Journal of Pharmaceutical Science and Technology Advanced Virus Detection Technologies Interest Group (AVDTIG): Efforts on High Throughput Sequencing (HTS) for
Recommendations for the introduction of metagenomic next-generation sequencing in clinical virology, part II: bioinformatic analysis and reporting Jute J.C. de Vira ¹¹ , Juliane R. Brown ¹ , Natacha Couto, ¹ Martin Bert ²¹ , Philippe I Mercier ¹ , gor Sidour ¹ , Arau Sayine ¹¹ , Mario Hosemani, ¹ Alba Perez-Catalunia ¹² , Ellen C. Cubo', Claudia Bachoni, ¹ Auba Kalodd, ¹ Domis Schnitz ¹ , Kurtina Tiska, ¹ Sober Mathumati, Dark Höper ¹ , Marta Hernande ¹¹ , ¹ Easheth Ponlhamme, Stockl ¹ , Anam Lehrand ¹ , Marta Hernande ¹¹ , ¹ Easheth Ponlhamme, Stockl ¹ , Anam Lehrand ¹ , Marta Hernande ¹¹ , ¹ Easheth Ponlhamme, Stockl ¹ , Anam Lehrand ¹ , Marta Hernande ¹¹ , ¹ Easheth Ponlhamme, Stockl ¹ , Anam Lehrand ¹ , Marta Hernande ¹¹ , ¹ Easheth Ponlhamme, Stockl ¹ , Anam Lehrand ¹ , ¹ Okarda Marta Hernande ¹¹ , ¹ Easheth Ponlhamme, Stockl ¹ , ¹ Anam Lehrand ¹ , ¹ Okarda Ponlhamme, ¹ Okarda Ponlhamd ¹ , ¹ Okarda Ponlhamme, ¹ Okarda Ponlhame, ¹ Okarda Ponlhamme, ¹ Okarda Ponlhame, ¹ Okarda Ponlhamme, ¹ Okarda Ponlhamme, ¹ Okarda Ponlhamme, ¹ Okarda Ponlhame, ¹ Okarda Ponlhamme, ¹ Okarda Ponlhame, ¹ Oka	Recommendations for the introduction of metagenomic high-throughput sequencing in clinical virology, part E. Wer lab procedure F. Aviera Löges Laborato ⁺¹ , Johlmer Renov, Niede Froncedure Sander Van Bohernen, Ontrof Clinick, Arza Sayiner, Tima Varehan Madem, Eera Auvinen, Verens Kufer, "Muhai Huber, "Christopher Roftguer," March 200ges ⁻¹ , Match Moren, A. Fedra March Huber, "Christopher Roftguer, "March 200ges ⁻¹ , March Hernandez", Reland Wolenkamp, J. Las ond Erleber, "Bok Schummar, Natech Octoor," Kanoline Learinger," Patter Simmonds, "March Berer, "Dirk Höger," Sergie Kannings,", "Endone Learinger, "Peres Simmonds," March 100ger, "March 200g, "Sergie Kannings,", Kanoline Learinger, "Sergie Kannings, "March 200g, "Sergie Kannings,", "Endone Learinger, "Sergie C. Marce, "Area Nate, "Andre Berer," End, "Sergie Kannings,", Sergie C. Marce, "Sergie K. Marce, Marce Marce, "Sergie Kannings,", "Sergie Kannings,", "Sergie Kannings, "Sergie Kannings,", "Sergie Kannings,", "Sergie Kannings,", "Sergie Kannings, Sergie Kannings	Virus Detection Arts: Stan, Dominick A. Vacarte, Jean-Pol Cassart, et al. PDA J Pharm 25 and Tred: 2017, 20 591,595 Access the most incert version at dox 10.57316cdappt.2016.007161
Perspective Current Perspectives on High-Throughput Sequencing (HTS) for Adventitious Virus Det Upstream Sample Processing and Library Prej Siemon H. Ng ¹⁴ , Cassandra Braxton ² , Marc Eloit ³⁴ , Szi Fei Feng ³ , Romain Fag Laurent Mallet ⁷ , Edward T. Mee ³ , Sarmitha Sathiamoorthy ¹⁴ , Olivier Vandeputte Arifa S. Khan ¹⁰	tection: paration Considerations for Optimization of High-Through Sequencing Bioinformatics Pipelines for View Determine	Paul W. Barone, Michael E. Wichel, James C. Leurg, Ham T. M. Hussein, Flora J. Keumuriani, James Bourssa ^{1,4} , Audrey Stussi ^{10,4} , Dyou C. Horry, ¹ Micy Cheng, ¹ Houman Delphanina ^{10,4} Lionel Gerentes ¹ , James Gillert ^{4,5} , Dan Gold', Robert Kiss ^{10,6} , Thomas R. Keill ¹ , Rend Labatet ¹ , Yuling L ^{10,4} , Jürgen Müllberg ¹¹ , Laurent Müllet ^{10,4} , Christian Menzel ¹¹ , Mark Moody ^{10,4} , Serge Monpoeho ¹⁴ , Mark Murphy ¹⁴ , Mark Parsiet ^{11,4} , Nathan J. Andr ¹⁴ , David Roush ¹⁴ , ¹⁴
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Reference Standards and Database





Creation a Ph. Eur General Chapter

- Expert group 15: "Human Vaccines and Sera" is working towards a new Ph. Eur. general chapter on "High Throughput Sequencing for the detection of extraneous agents (2.6.41)"
- This chapter will provide details on important considerations when implementing a HTS for adventitious virus detection assay and provide guidance on the validation of such a method.

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• Support the implementation the HTS technology in a GMP environment

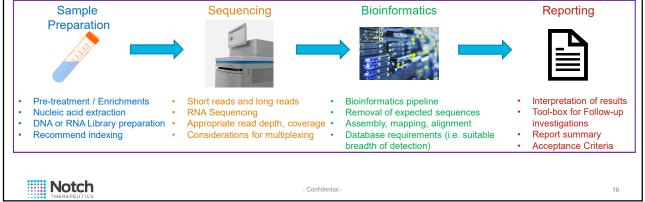
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Describe different HTS technology and the multiple possible approaches

• Provide a standardization of critical components that are part of the assay

Topics to be covered:

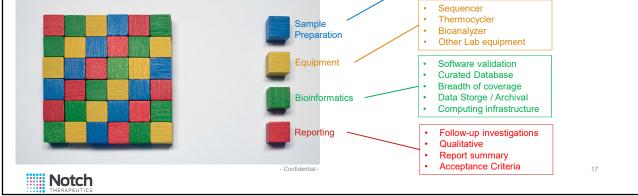
• Different analysis (Genomics, Transcriptomics, Targeted) and sample types (Cell banks, Viral seeds, Harvest)



Validation Approach

· Recommends a modular approach where individual components can be qualified / validated separately before validating the entire method





Opportunity to streamline the method validation

- Validation of a complex method and platform that doesn't necessarily follow ICH Q2
 - Breadth of detection (specificity) demonstrated using a few model viruses representing different types of viruses
 - Minimum characterization requirements for model viruses
 - Propose to minimized the number of replicates to two or three runs Achieve a balance between cost and demonstrating consistency
 How to demonstrate the equivalency of the results between replicates?
 - Validation parameters
 - Demonstrating reproducibility for qualitative vs quantitative assays
 Sensitivity and LOD
- Clarification that a head-to-head comparison is not needed to replace in vivo tests



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Promote Standardization for the use of HTS	
 Provide the first strategic frame-work for the HTS validation approach 	
Clarify regulatory expectation	
 Promote the use of the HTS technology in a GMP environment 	
 Potentially serve as a foundation for evaluating HTS dossier submission 	
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Expansion of HTS applications



Expansion of HTS appli	cations
	Biotechnology
 Gene and Cell Therapy 	Systems & Synthetic Biology · Nanobiotech · Medicine Journal Research Article
Published online 8 December 2020 Nucleic Acids Research, 2021, Vol. 49, No. 3 e16 doi: 10.1093/narigkou1152	Enhanced CHO Clone Screening: Application of Targeted Locus Amplification and Next-Generation Sequencing Technologies for Celluica Development
Genome-wide integration site detection using Cas9 enriched amplification-free long-range sequencing	Cell Line Development Samuel H. Aeschlimann. Christian Graf. Dmytro Mayilo. Hélène Lindecker, Lorena Urda. Nora Kappes. Alicia Leone Burr, Marieke Simonis. Erik Splinter, Max van Min. Holger Laux 🕱
Joost van Haasteren [†] , Altar M. Munis ^{©†} , Deborah R. Gill and Stephen C. Hyde	First published: 22 February 2019 https://doi.org/10.1002/biot.201800371 Citations: 4
Gene Medicine Group, Nuffield Division of Clinical Laboratory Sciences, Radcliffe Department of Medicine, University of Oxford, Oxford, UK	> Nat Biotechnol. 2014 Oct;32(10):1019-25. doi: 10.1038/nbt.2959. Epub 2014 Aug 17.
BIOTECHNOLOGY	Targeted sequencing by proximity ligation for comprehensive variant detection and local haplotyping
Multiplexed clonality verification of cell lines for protein biologic production	Paula J P de Vree ¹¹ , Elzo de Wit ² , Mehmet Yilmaz ³ , Monique van de Heijning ³ , Petra Klous ³ , Marjon J A M Verstegen ⁴ , Vi Wan ⁴ , Hans Teunissen ⁴ , Peter H L Krijger ⁴ , Geert Geeven ⁴ , Paul P Eilk ⁵ , Daoud Sie ⁵ , Bauke Yista ⁵ , Lorette O M Hulsman ⁶ , Marieke F van Dooren ⁶ ,
Sone A. Ubiter, Juni Una, Yalu Wolge Weishou Hulge First published: 07 February 2020 https://	Laura J C M van Zutve Marion Cornelissen ⁸
Article Published: 15 June 2020 CHANGE-seq reveals genetic and epigenetic of CRISPR-Cas9 genome-wide activity Circera R. Larzenton. Nikolay. L. Mainin Sitchard (Zhang, Yang Gatiyan L. Yanghua He Xin Lan Kazey, Lieden Maran Kata Natala G. Kolmakova Christopher L. Ergheni Sitchcov, Sammath Madagh, Gieder Arencia, Band Mayo, Cheng & Shenge	Les Elsanor Cowley? Peternen Glan Gla Peternen Glan Gla
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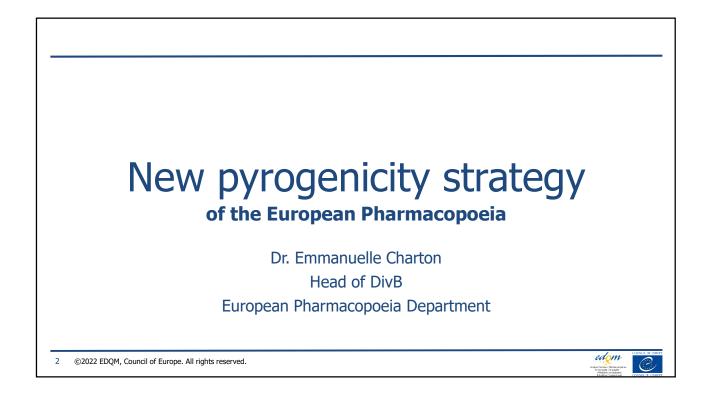
Summary and Conclusion

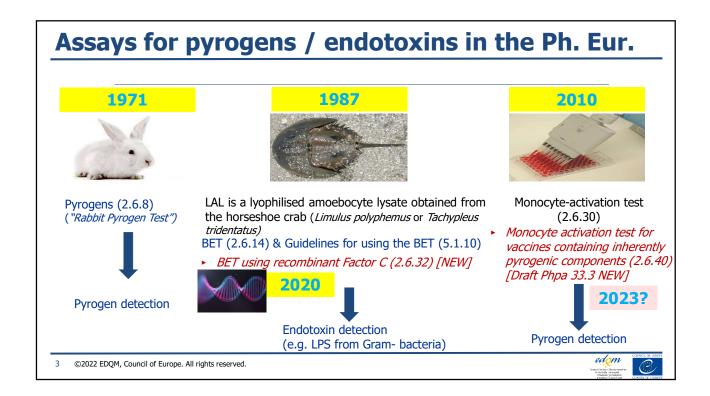
- HTS is a technology capable of known and unknown viral adventitious agent in a biological samples.
- Recent achievements and alignment within the community towards the implementation of a viral adventitious agent detection by HTS as a GMP specification test.
- Drafting of a new Ph. Eur. general chapter will provide a suitable approach towards validation of the HTS assay and promote wider adaption of the HTS technology

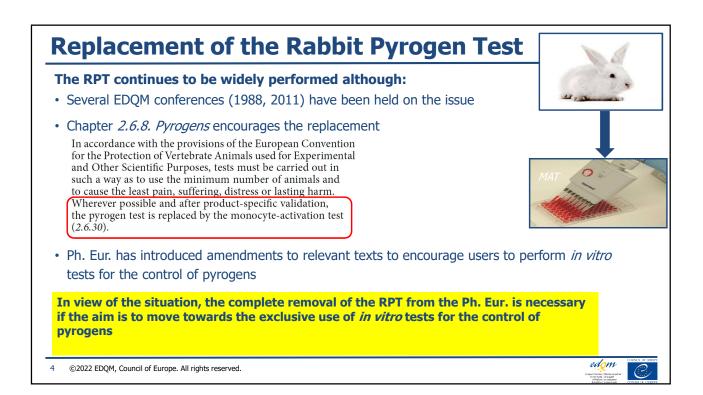
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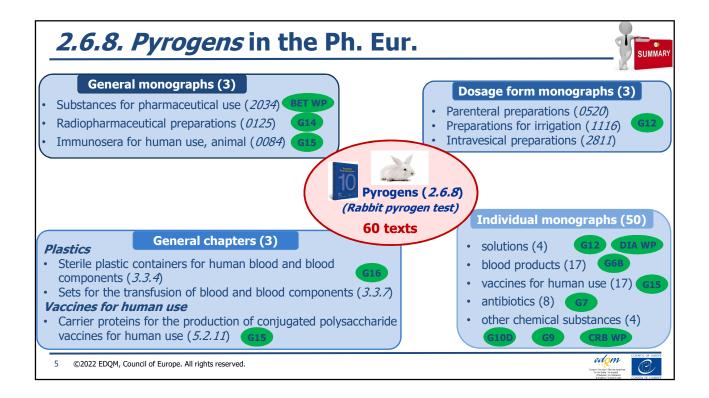
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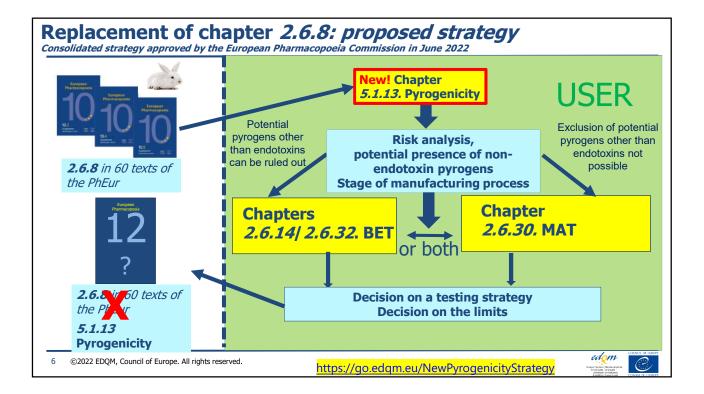


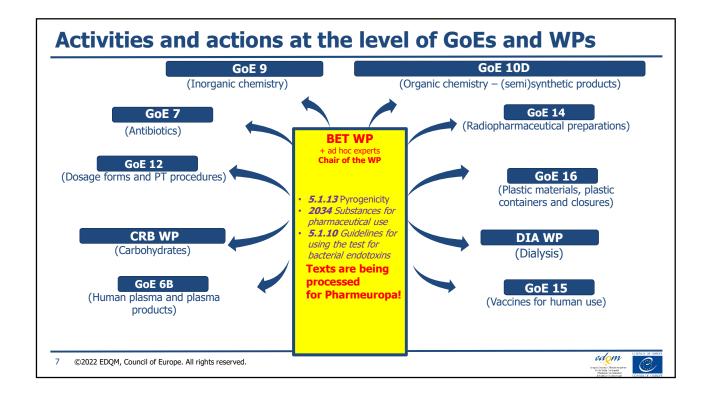


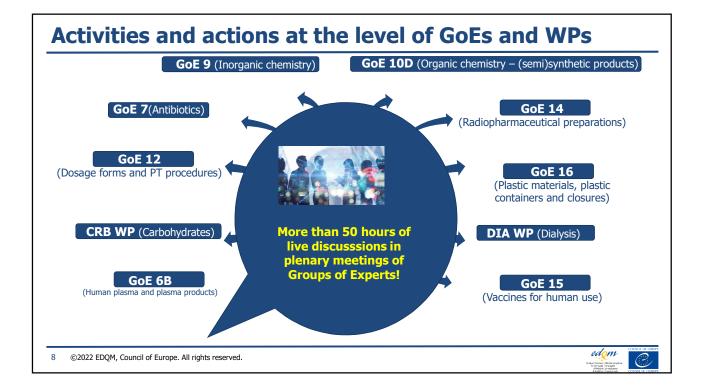


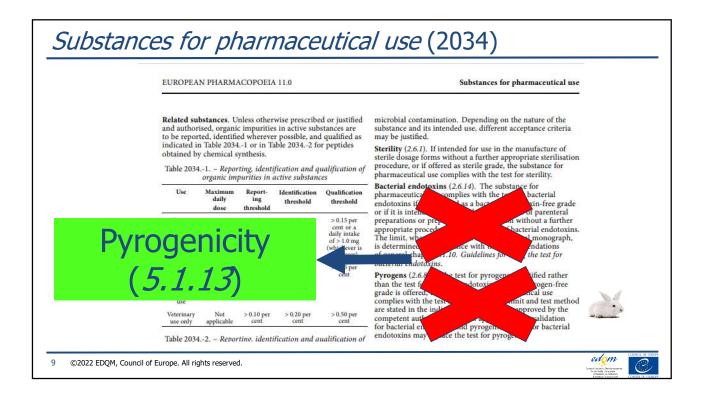


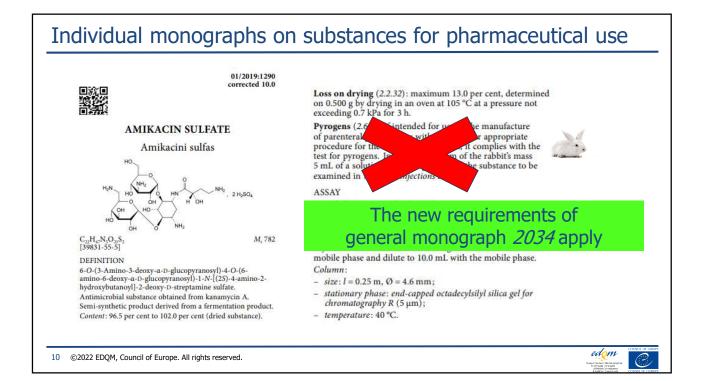


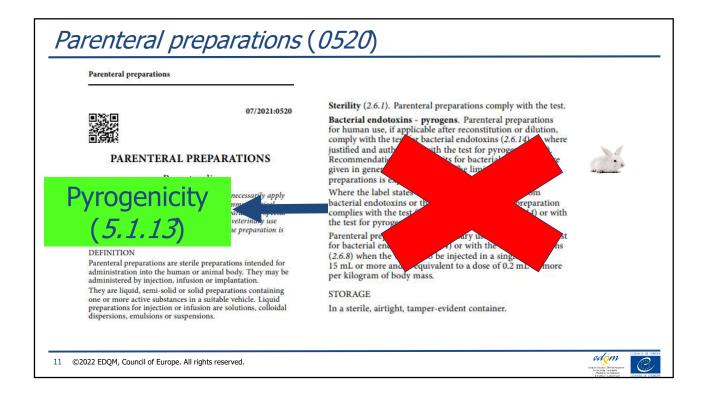






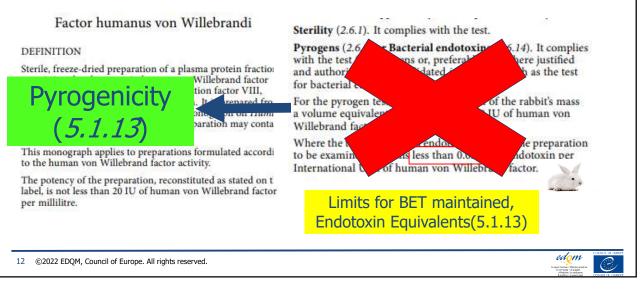






Plasma-derived products

HUMAN VON WILLEBRAND FACTOR



Monograph/chapter		Requirement for RPT		
Hepatitis B- containing vaccines	- Нер В (1056) - DT-Нер В* (2062) - DTaP-Нер В* (1933)	- RPT on the final lot		
3-O-Desacyl-4'-mono	phosphoryl lipid A (MPL) (2537)	- RPT on an intermediate		
Haemophilus influenza type b- containing vaccines	- Hib (1219) - DTaP-Hib (1932) - DTaP-IPV-Hib (2065) - DTwP-IPV-Hib* (2066)	 - RPT as a process validation requirement - RPT on the final lot if any vaccine component prevents the determination of endotoxin 	[
	- DTaP-IPV-Hep B-Hib (2067) - Hib-Men C (2622)	- RPT as a requirement during product development	+ Revise general monograph <i>Vaccines</i>	
Meningococcal	- Men PS vaccine (0250)	- RPT on an intermediate and on the final lot	for human use (0153)	
vaccines	- Men C conjugate vaccine (2112) - Men A, C, W135, Y conjugate vaccine (3066)	- RPT as a process validation requirement		
Pneumococcal vaccines	- Pneumococcal polysaccharide vaccine (0966)	- RPT on final lot		
	- Pneumococcal conjugate vaccine (2150)	- RPT as a requirement during product development		
Rabies vaccine (0216)		- RPT on the final lot in case non-endotoxin pyrogens are present	*monographs will be suppressed from the Ph. Eur. as of July 2023	
Tick-borne encephali	tis vaccine (1375)	- RPT on the final lot	(Supplement 11.2)	
Carrier proteins for tl (5.2.11)	ne production of conjugated vaccines	- RPT for <i>N. meningitidis</i> outer membrane protein complex (OMP)	Long to Table Store store the	

General monograph Vaccines for human use (0153)

	PRODUCTION
NOTE ON THE GENERAL MONOGRAPH Pyrogenicity . The section on Bacterial endotoxins in the Tests part of the monograph has been replaced with a new section on Pyrogenicity, referring to new general chapter 5.1.13 Pyrogenicity which provides guidance for selection and implementation of a suita- ble test for pyrogenicity (test for bacterial endotoxins or monocyte-activation test). In addition, a statement has been introduced under General provisions in the Production part of the monograph to stress the need to characterise pyrogenicity during development studies and whenever revalidation is necessary. This revision of general monograph 0153 is part of a broader exercise affecting multiple Ph. Eur. tests and aiming at the complete suppression of the rabbit pyrogen test from the Ph. Eur. As part of this exercise, the following texts have been published in the same issue of Phar- meuropa: 1) new general chapter 5.1.13 Pyrogenicity; 2) monographs on individual vac- cines for human use that were revised to delete the reference to th. The revised individual monographs no longer contain any mentio and, as a result, the requirements of general monograph 0153 for General provisions and Tests) will apply.	 General provisions. The production method for a given product must have been shown to yield consistently batches comparable with the batch of proven clinical efficacy, immunogenicity and safety in man. Product specifications including in-process testing should be set. Specific requirements for production including in-process testing are included in individual monographs. Where justified and authorised, certain tests may be omitted where it can be demonstrated, for example by validation studies, that the production process consistently ensures compliance with the test. Unless otherwise justified and authorised, vaccines are produced using a seed-lot system. The methods of preparation are designed to maintain adequate immunogenic properties, to render the prenaration harmless and to prevent contamination with extraneous agents. Pyrogenicity is characterised during development studies and controlled whenever revalidation is necessary. Guidance for selection of a suitable pyrogenicity test is given in general chapter <i>5.1.13</i>. TESTS Vaccines comply with the tests prescribed in individual monographs including, where applicable, the following:
Importantly, the revision of the monograph does not call into question established manufacturers' strategies to control the pyrogenicity of their products using the test for bacterial endotoxins that were authorised by the competent authority, and is not intended to prompt a retrospective assessment on pyrogenicity.	Bacterial endotoxins. Unless otherwise justified and authorised, a test for bacterial endotoxins is carried out on the final product. Where no limit is specified in the individual monograph, the content-of-bacterial endotoxins determined by a suitable method (2.6.14) is less than the limit approved for the particular-product. Pyrogenicity. The vaccine complies with a suitable test for pyrogenicity. Guidance for selection of a test is given in general chapter 5.1.13. Where no limit is specified in the individual monograph, it complies with the limit approved for the particular product.
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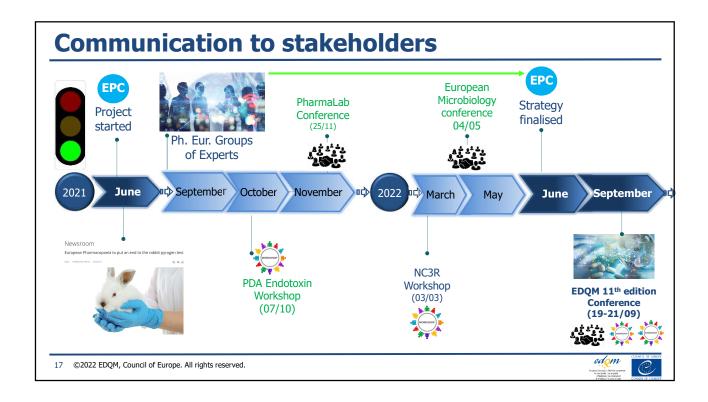
NOTES ON THE TEXTS "It should be noted that the exercise will ultimately lead to the suppression of general chapter 2.6.8 from the Ph. Eur. Manufacturers still using the rabbit pyrogen test are strongly encouraged to take the necessary steps to proceed with its replacement by a suitable in vitro alternative (e.g. the monocyte-activation test), in line with the new

requirements of this general monograph."

• "Importantly, the revision of this text does not call into question strategies involving the test for bacterial endotoxins that are already used by manufacturers to control the pyrogenicity of their products and have been authorised by the competent authority, nor is it intended to prompt a retrospective assessment of pyrogenicity."

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WHAT	WHO	٧	WHEN
		Publication in PhPa	Envisaged implementatio date
(5.1.13) (together with revision of 5.1.10)	BET WP	0	
REVISION			
2.6.30	BET WP	\bigcirc	
2034	BET WP	<u> </u>	
0520	G12 with BET WP support	0	
remaining texts	GoE/WP with BET WP support	<u> </u>	
DELETE Pyrogens (2.6.8)			
		April	July July



EPAA/EDQM International Public Conference

To mark the first official milestone of the strategy, i.e. the publication of revised Ph. Eur. texts omitting the RPT in Pharmeuropa 35.1 (January 2023)



Date: 14-16 February 2023

Venue: European Commission premises, Brussels

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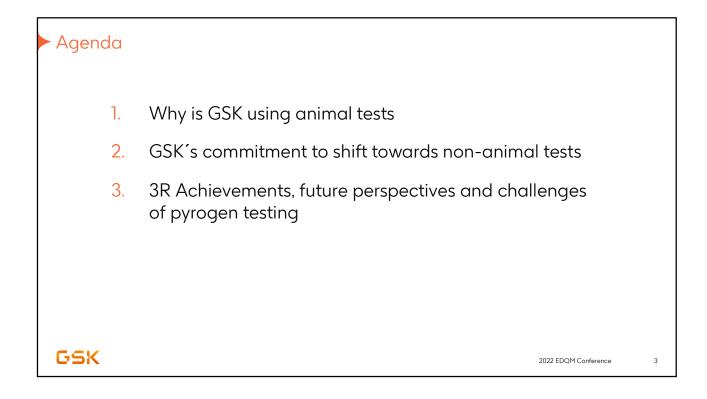
► Disclosure

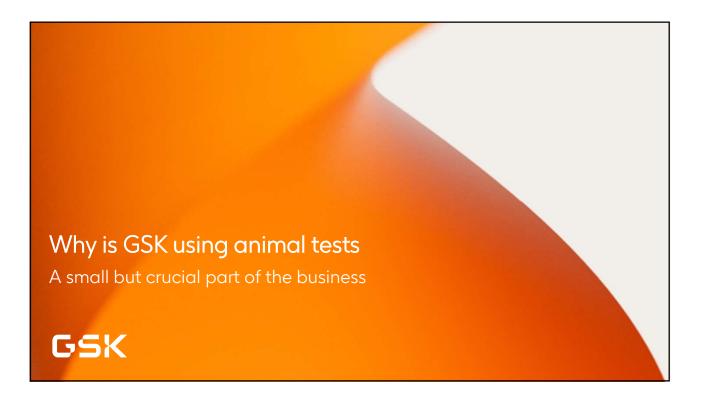
Shahjahan Shaid is an employee of the GSK group of companies. This work was sponsored by GlaxoSmithKline Biologicals SA.

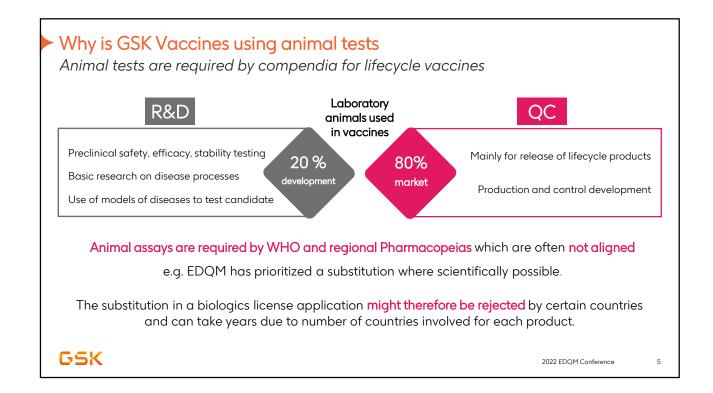
GSK

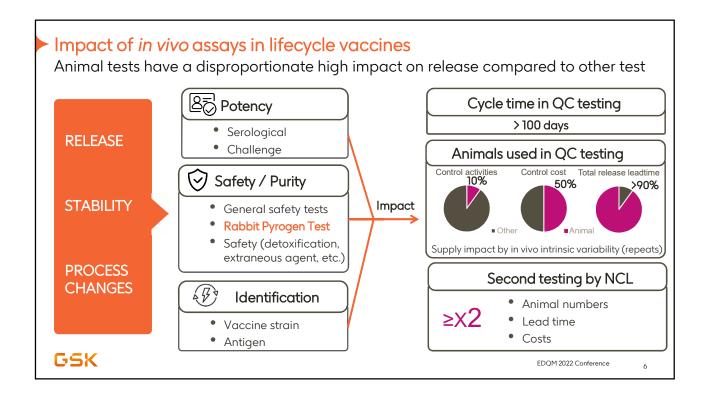
2022 EDQM Conference

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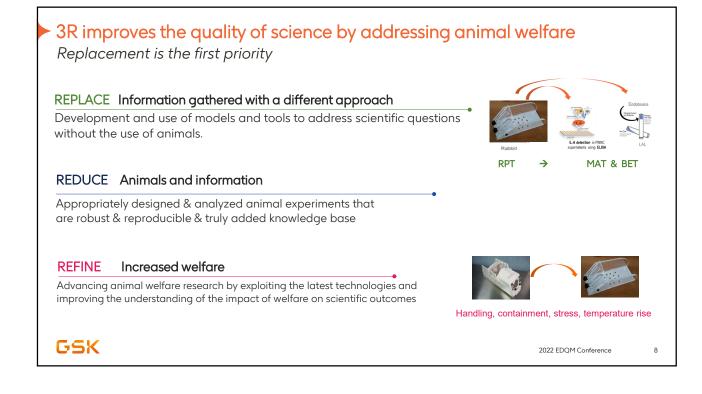


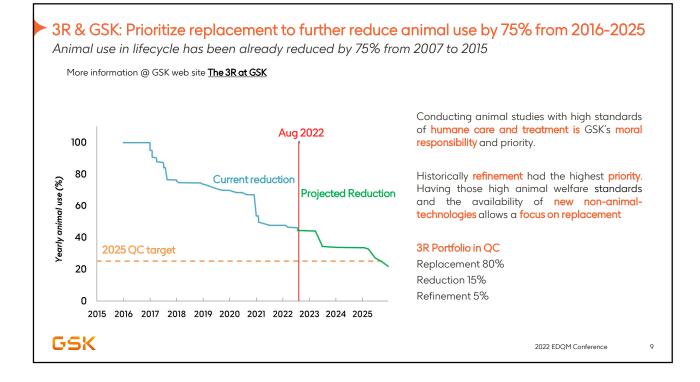


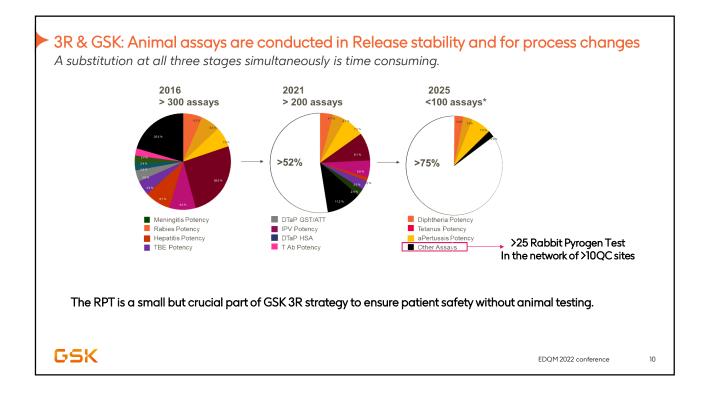
GSK's Vaccine commitment to shift towards non-animal tests

3R strategy: created to respond to the changing environment

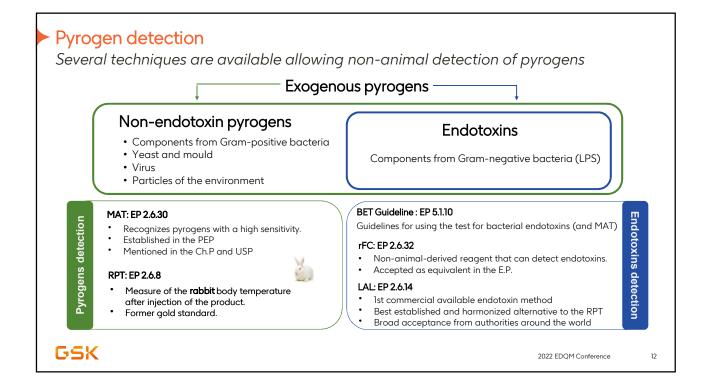
GSK

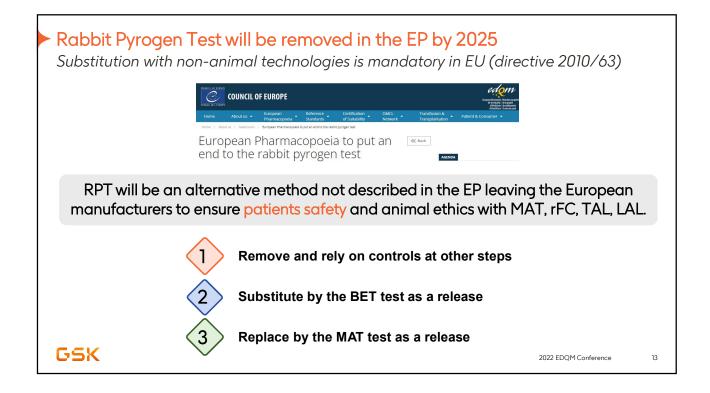


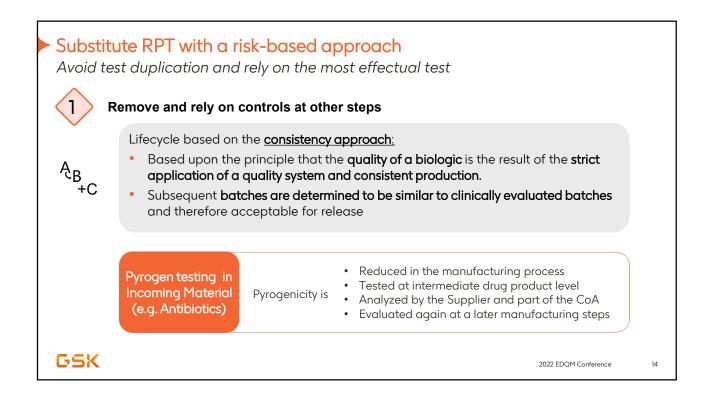


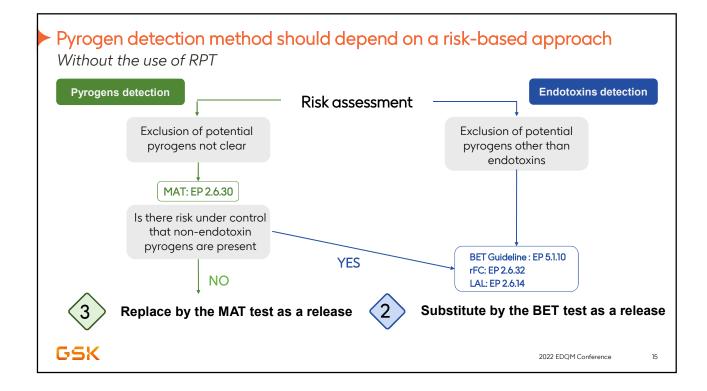


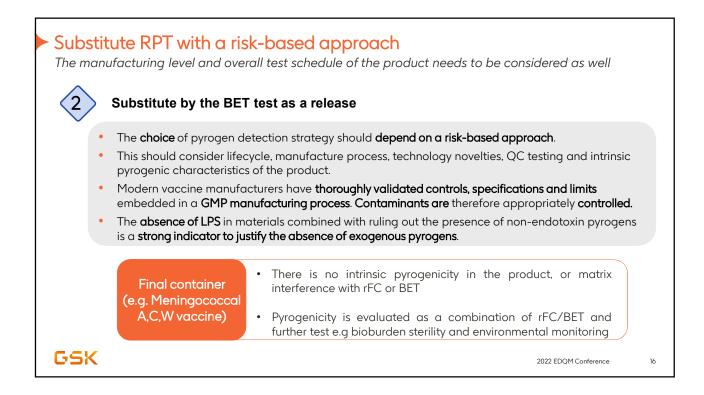


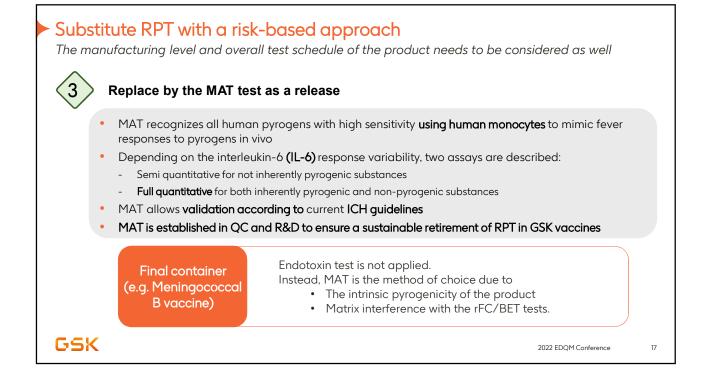


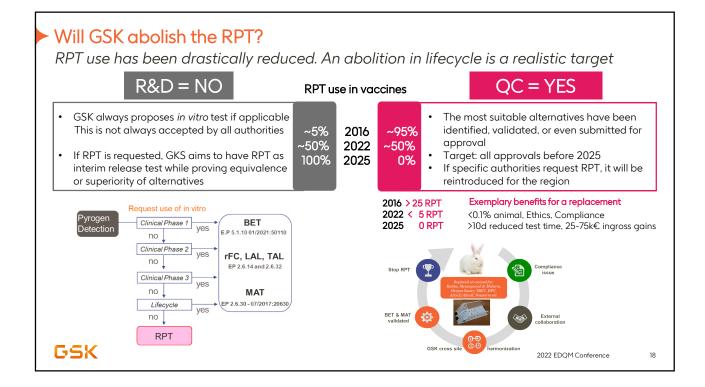












To avoid the use of RPT beyond lifecycle, harmonization is required RPT is still considered as gold standard in certain regions beyond Europe, impacting abolition at R&D level While the BET is preferred by the user and widely accepted as a substitution method by Health authorities it is not always applicable.				
-	The regula	tory landscape beyond Europe regarding acceptance of MAT		
	WHO	Only states RPT (proposal has been drafted)		
	USA	USP 151 – Validated and equivalent in vitro test may be used in place of RPT where appropriate		
	China	MAT is included to be conducted in addition to RPT for guideline 9301: "Application of Safety Tests for Injection"		
Rep. of Korea		No mention of MAT making it an alternative method.		
	Japan	No mention of MAT making it an alternative method . Requires proving superiority in terms of accuracy and precision		
	India	Considers MAT a suitable alternative test requires to show equivalence		
GSK	Brazil	Pyrogen test required. Expected to add MAT in their Pharmacopoeia 2022 EDQM Conference 19		

